

SEQ ID NO:1

1 CGTCACCACCA ~~GTA~~GGTCCCT CGGCTCAGTC GGCCCAGCCC CTCTCAGTCC
51 TCCCCAACCC CCACAACCGC CGCGGGCTCT GAGACGGGGC CCGGGCGGGC
101 CGGGCAGCAG CTGCAGCATC ATCTCACCC TCCAGCCATG GAAGACCTGG
151 ACCAGTCTCC TCTGGTCTCG TCCCTGGACA GCCCACCCCG GCCGCAGCCC
201 GCGTTCAAGT ACCAGTTCGT GAGGGAGCCC GAGGACGAGG AGGAAGAAGA
251 GGAGGAGGAA GAGGAGGACG AGGACGAAGA CCTGGAGGAG CTGGAGGTGC
301 TGGAGAGGAA GCCCGCGGCC GGGCTGTCCG CGGCCCCAGT GCCCACCGCC
351 CCTGGCCGCCG CGCGCGCCCT GATGGACTTC GGAAATGACT TCGTGCCGCC
401 GCGGCCCCGG GGACCCCTGC CGGGCGCTCC CCCCGTCGCC CGGGAGCGGC
451 AGCCGTCTTG GGACCCGAGC CGGGTGTCTG CGACCGTGCC CGGCCATCC
501 CGCTGTCTG CTGCCGCAGT CTGCCCTCC AAGCTCCCTG AGGACGACGA
551 GCCTCCGGCC CGGCCTCCCC CTCCCTCCCC CGCCAGCGTG AGCCCCCAGG
601 CAGAGCCCGT GTGGACCCCG CCAGCCCCGG CTCCCGCCGC GCCCCCTCC
651 ACCCCGGCCG CGCCCAAGCG CAGGGCTCC TCAGGGCTCAG TGGTTGTTGA
701 CCTCCTGTAC TGGAGAGACA TTAAGAAGAC TGGAGTGGTG TTTGGTGCCA
751 GCCTATTCCCT GCTGCTTTCA TTGACAGTAT TCAGCATTGT GAGCGTAACA
801 GCCTACATTG CCTTGGCCCT GCTCTCTGTG ACCATCAGCT TTAGGATATA
851 CAAGGGTGTG ATCCAAGCTA TCCAGAAATC AGATGAAGGC CACCCATTCA
901 GGGCATATCT GGAATCTGAA GTTGCTATAT CTGAGGAGTT GGTCAGAAG
951 TACAGTAATT CTGCTCTTGG TCATGTGAAC TGCACCGATAA AGGAACTCAG
1001 GCGCCTCTTC TTAGTTGATG ATTTAGTTGA TTCTCTGAAG TTTGCAGTGT
1051 TGATGTGGGT ATTTACCTAT GTTGGTGCCT TGTATAATGG TCTGACACTA
1101 CTGATTTGG CTCTCATTTC ACTCTTCAGT GTTCCCTGTTA TTTATGAACG
1151 GCATCAGGCA CAGATAGATC ATTATCTAGG ACTTGCAAAT AAGAATGTTA
1201 AAGATGCTAT GGCTAAAATC CAAGCAAAAA TCCCTGGATT GAAGGCCAAA
1251 GCTGAATQAA AACGCCAAA ATAATTAGTA GGAGTTCATC TTTAAAGGGG
1301 ATATTCAATT GATTATAACGG GGGAGGGTCA GGGAAAGAACG AACCTTGACG

SEQ ID NO. (con't)

1351 TTGCAGTGCA GTTTCACAGA TCGTTGTTAG ATCTTTATTT TTAGCCATGC
1401 ACTGTTGTGA GGAAAAATTAA CCTGTCTTGA CTGCCATGTG TTCATCATCT
1451 TAAGTATTGT AAGCTGCTAT GTATGGATT AAACCGTAAT CATACTTTT
1501 TCCTATCTGA GCCACTGGTG GAATAAAAAA CCTGTATATT TTACTTTGTT
1551 GCAGATAGTC TTGGCGCATC TTGGCAAGTT GCAGAGATGG TGGACCTAGA
1601 AAAAAAAA AAAAAGCCCT TTTCAGTTTG TGCACTGTGT ATGGTCCGTG
1651 TAGATTGATG CAGATTTCT GAAATGAAAT GTTTGTAG ACAGAGATCAT
1701 ACCGGTAAAG CAGGAATGAC AAAGCTTGCT TTCTGGTAT GTCTAGGTG
1751 TATTGTGACT TTACTGTTA TATTAATTGC CAATATACTT AAATATAGAT
1801 TATAATGTA TAGTGTTC CAACAGCTTAG ACCTTTACCT TCCAGCCACC
1851 CCACAGTGCT TGATATTTCA GAGTCAGTCA TTGGTTATAC ATGTGTAGTT
1901 CCAAAGCACA TAAGCTAGAA GAAGAAATAT TTCTAGGAGC ACTACCACCT
1951 GTTTCAACA TGAAATGCCA CACACATAGA ACTCCAACAA CATCAATTTC
2001 ATTGCACAGA CTGACTGTAG TTAATTTGT CACAGAATCT ATGGACTGAA
2051 TCTAATGCTT CCAAAATGT TGTGTTGTTG CAAATATCAA ACATTGTTAT
2101 GCAAGAAATT ATTAATTACA AAATGAAGAT TTATACCATT GTGGTTAAG
2151 CTGACTGAA CTAAATCTGT GGAATGCATT GTGAACTGTA AAAGCAAAGT
2201 ATCAATAAAG CTTATAGACT TAAAAAAA AAAAAAAA

SEQ ID NO:2

1 MEDLDQSPLV SSSDSPPRPQ PAPKTYQFVRE PEDZEEEEE EEEDEDEDLE
51 ELEVLERKPA AGLSAAPVPT APAAGAPLMD FGNDPVFFAF RGFLPAAPPV
101 APERQPSWDP SVVSSTVPAP SPLSAAAVSP SKLPEDDEPP ARPPPPPAS
151 VSPQAEPVWT PPAPAPAAPP STPAAPKRRG SSG3VVVYDLL YWRDIKKTGV
201 VFGASLFLLL SLTVFSIVSV TAYIALALLS VTISPRIYKG VIQAIQKSDE
251 GHPFRAYLES EVAISEELLYQ KYSNSALGHV NCTIKELRRL PLVDDLVDSL
301 KFAVLNNWPT YVGALPNGLT LLILALISLF SVFVIYERHQ AQIDHILGLA
351 NKNVKDAMAK IQAKIPGLKR KAE